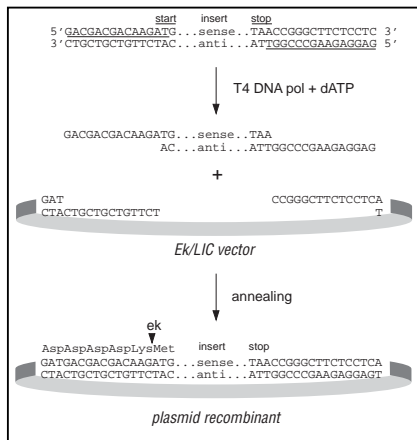


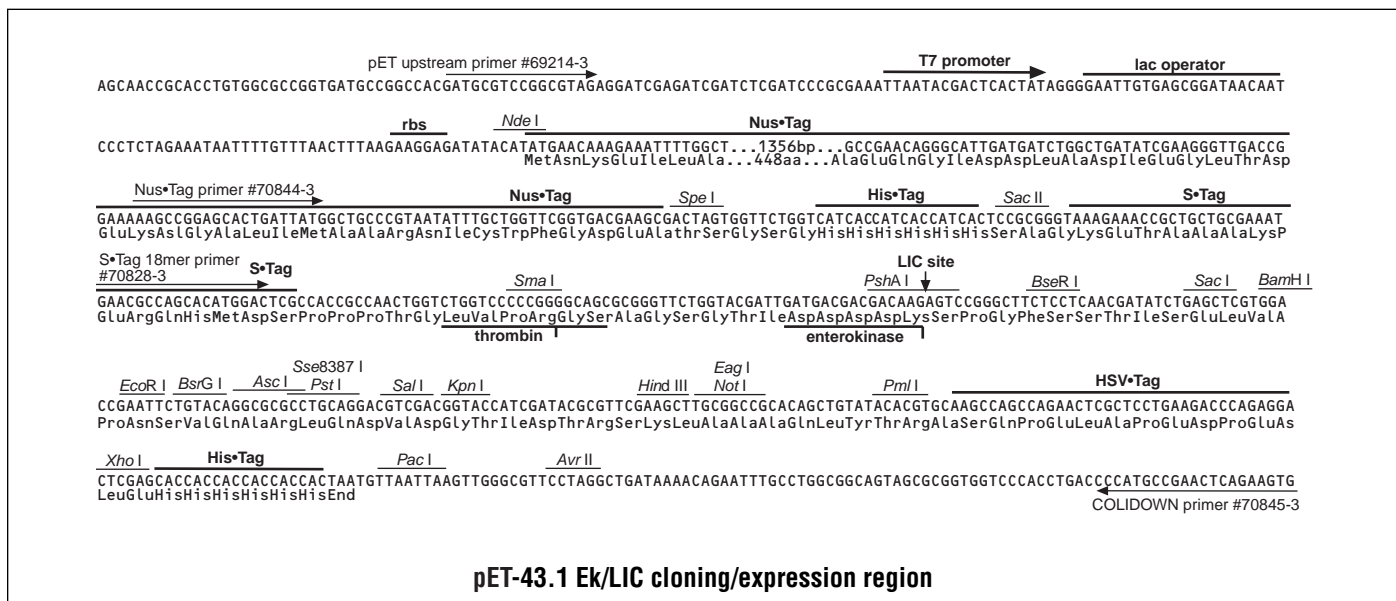
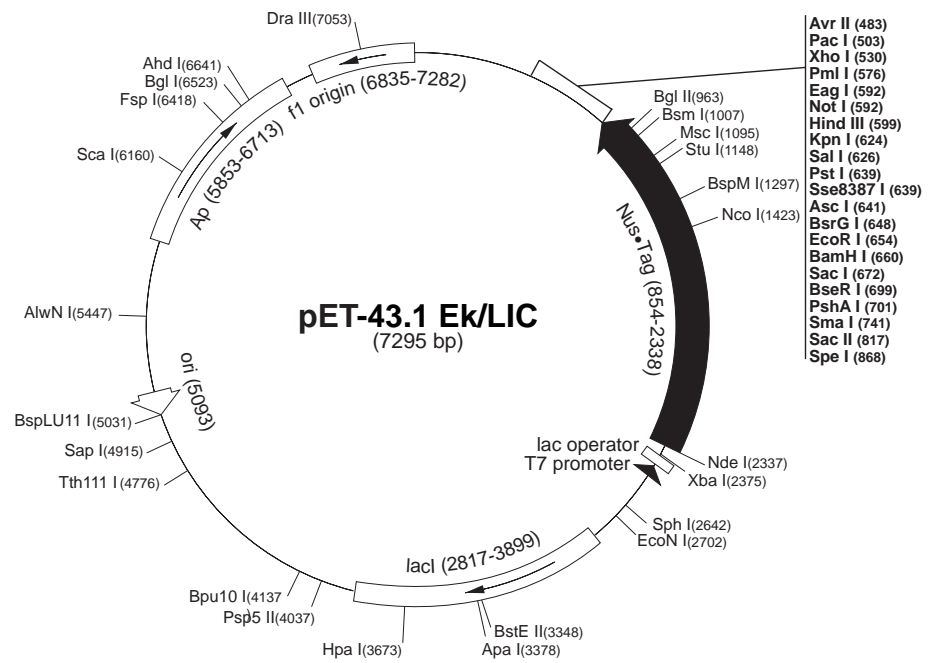
pET-43.1 Ek/LIC Vector

pET-43.1 EK/LIC sequence landmarks

T7 promoter	2410-2426
T7 transcription start	2409
Nus•Tag coding sequence	854-2348
His•Tag coding sequence	821-838
S•Tag coding sequence	767-811
Multiple cloning sites (<i>BseR I-Xho I</i>)	699-530
HSV•Tag coding sequence	536-571
His•Tag coding sequence	512-529
T7 terminator	26-72
<i>lacI</i> coding sequence	2817-3899
pBR322 origin	5093
<i>bla</i> (Ap) coding sequence	5854-6714
f1 origin	6835-7282



The pET-43.1 Ek/LIC vector is prepared for rapid, directional cloning of PCR-amplified DNA for high-level expression of polypeptides fused with N-terminal Nus•Tag™, His•Tag® and S•Tag™ sequences. Using specifically designed primers for amplification and the pET-43.1 Ek/LIC Vector Kit (Cat. No. 71072-3), inserts can be efficiently cloned without the need for restriction digestion or ligation. Unique sites are shown on the circle map. Note that the sequence is numbered by the pBR322 convention, so the T7 expression region is reversed on the circle map. The cloning/expression region of the coding strand transcribed by T7 RNA polymerase is shown below. The f1 origin is oriented so that infection with helper phage will produce virions containing single stranded DNA that corresponds to the coding strand. Therefore, single stranded sequencing should be performed using the COLIDOWN primer (Cat. No. 70845-3). Vector encoded sequence can be completely removed when cloning into the Ek/LIC site (as shown below left) by cleaving the fusion protein with enterokinase.



pET-43.1 Ek/LIC cloning/expression region

pET-43.1 Ek/LIC Restriction Sites

Enzyme	# Sites	Locations	Enzyme	# Sites	Locations	Enzyme	# Sites	Locations				
AatII	2	633 2208	DdeI	15		Sau3AI	42					
AccI	4	580 627 1549 4801	DpnI	42		Sau96I	20					
AcII	99		DraI	3	6063 6755 6774	Scal	1	6160				
AflIII	4	575 608 3167 5031	DraIII	1	7053	ScrFI	33					
AhdI	1	6641	DrdI	3	4724 5139 7008	SfaNI	23					
AluI	27		DsaI	4	814 984 1423 2604	SfcI	7	635 2314 2409 5296 5487				
Alw26I	8	371 2864 3269 3395 3782	EaeI	8	592 991 1093 1380 2475			6395 7272				
		4672 5798 6575			2607 3841 6248	SgrAI	2	1919 2486				
AlwI	20		EagI	1	592	SmaI	1	741				
AlwNI	1	5447	EarI	3	2785 4915 5842	SpeI	1	848				
Apal	1	3378	Ecl136II	1	670	SphI	1	2642				
ApalI	4	3147 4845 5345 5969	Eco47III	2	2572 4285	Sse8387I	1	639				
ApoI	7	466 654 788 2323 3442	Eco57I	4	532 1846 5579 5975	SspI	2	877 6845				
		6855 6866	EcoNI	1	2702	StuI	1	1148				
Ascl	1	641	EcoO109I	3	53 2600 4037	StyI	4	57 483 1423 2119				
Aval	2	530 739	EcoRI	1	654	TaiI	25					
Avall	7	440 745 3719 4037 4316	EcoRII	13		TaqI	28					
		6277 6499	EcoRV	3	677 931 1522	TfiI	4	3846 4081 4585 5006				
AvrII	1	483	EheI	4	2491 2512 2626 3808	ThaI	52					
BamHI	1	660	FauI	20		TseI	28					
BanI	10	620 1470 2489 2510 2624	Fnu4HI	52		Tsp45I	10	859 1745 2132 3348 4470				
		3087 3806 3936 6688 7090	FokI	14				4683 4778 6168 6379 7226				
BanII	5	672 2551 2565 3378 7128	FspI	1	6418	Tsp509I	23					
BbsI	5	538 1228 3313 3652 4149	HaeI	15		TspRI	16					
BbvI	28		HaeIII	31		Tth111I	1	4776				
BcgI	3	3493 4608 6103	HgaI	15		VspI	4	2424 3852 3911 6466				
BclI	2	1953 3181	HhaI	59		XbaI	1	2375				
Bfal	10	70 484 849 2376 4010	HincII	4	628 920 1315 3673	XcmI	3	3023 3539 3557				
		4045 5526 6448 6783 7204	HindIII	1	599	XhoI	1	530				
BglI	1	6523	HinfI	16		XmnI	3	2271 4589 6041				
BglII	1	963	HpaI	1	3673							
BpmI	4	3005 3494 4558 6572	HphI	29		Enzymes that do not cut pET-43.1 Ek/LIC:						
Bpu10I	1	4137	KpnI	1	624	AflIII	Bsu36I	FseI	MunI	NheI	NsiI	PinAI
Bpu1102I	2	80 1298	MaeIII	22		PmeI	RsrII	SanDI	SexAI	SfiI	SgfI	SnaBI
BsaAI	3	576 4783 7053	MboII	25		SrfI	SunI	Swal				
BsaBI	3	2440 2450 4228	MluI	2	608 3167							
BsaHI	9	200 630 2205 2490 2511	MnlI	29								
		2625 3124 3807 6101	MscI	1	1095							
Bsal	2	371 6575	MseI	33								
BsaJI	17		MslI	11								
BsaWI	8	2 1596 3486 3989 4220	MspA1I	13								
		5237 5384 6345	MspI	42								
BseRI	1	699	MwoI	47								
BsgI	4	1976 3018 3218 4191	NarI	4	2490 2511 2625 3807							
BsiEI	10	595 1135 1273 1809 2190	NciI	20								
		3952 4947 5371 6123 6272	NcoI	1	1423							
BsiHKAI	11		NdeI	1	2337							
BsiI	26		NgoAIV	2	2477 7154							
BsmBI	2	3782 4672	NlaIII	28								
BsmFI	5	453 758 2628 4302 7268	NlaIV	25								
BsmI	1	1007	NotI	1	592							
Bsp1286I	15		NruI	3	1230 1655 2279							
BspEI	2	2 4220	NspI	4	2642 4376 4668 5035							
BspLU11I	1	5031	NspV	2	604 2265							
BspMI	1	1297	Pacl	1	503							
BsrBI	6	249 1343 2396 4964 5798	PfiMI	2	778 2749							
		7197	PleI	12								
BsrDI	6	1093 1474 3214 3580 6407	PmlI	1	576							
		6581	PshAI	1	701							
BsrFI	9	1047 1403 1467 1919 2477	Psp1406I	6	237 2829 4356 6039 6412							
		2486 2853 6556 7154			6838							
BsrGI	1	648	Psp5II	1	4037							
BsrI	29		PstI	1	639							
BssHII	5	641 1567 1759 2248 3578	PvuI	3	1273 2190 6272							
BssSI	4	665 1639 5204 5972	PvuII	4	586 3767 3860 4622							
Bst1107I	2	581 4802	RcaI	4	2565 5751 5800 5832							
BstEII	1	3348	RsaI	8	622 650 721 976 2198							
BstXI	3	2969 3098 3221			3314 4837 6160							
BstYI	13		SacI	1	672							
Cac8I	52		SacII	1	817							
Clal	5	615 1524 1602 2190 2444	Sall	1	626							
CviJI	113		SapI	1	4915							